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PATENT APPLICATION

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

1634
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Applicants: Arguello *et al.*

Appl. No.: 09/077,615

DEC 12 2002

Filing Date: October 23, 1998

TECH CENTER 1600/2900

Title: METHODS FOR SEPARATING AND/OR IDENTIFYING DNA MOLECULES

Docket No.: 028979-0103

Commissioner for Patents
Washington, D.C. 20231

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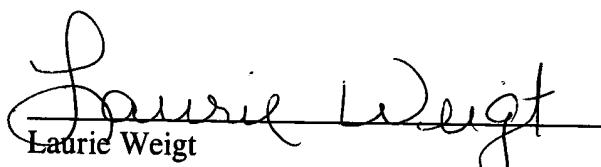
Date of Deposit: December 6, 2002

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- * Brief on Appeal (in triplicate)
- * Copies of Cited Thesaurus Pages (in triplicate)
- * Copies of Cited Dictionary Pages (in triplicate)
- * Copy of Cited Abstract (in triplicate)
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DEC 12 2002

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

TECH CENTER 1600/2900

Appellants: Rafael ARGUELLO *et al.*

Title: METHODS FOR SEPARATING AND/OR IDENTIFYING DNA MOLECULES

Appl. No.: 09/077,615

#35
10

Filing Date: 10/23/1998

Examiner: J. C. Einsmann

Art Unit: 1634

BRIEF ON APPEAL

Commissioner for Patents
Washington, D.C. 20231

Dear Sir:

This brief is in furtherance of the Notice of Appeal filed in this case on October 8, 2002. The required fee for filing an appeal brief, as set forth in 37 C.F.R. § 1.17(c) is included in our Check No. 730443. Any fee deficiency may be charged, or overpayment credited to, Deposit Account No. 50-2350. This brief is transmitted in triplicate and in conformance with the other requirements of 37 C.F.R. §1.192(a).

I. REAL PARTY IN INTEREST

Pel-Freez Clinical Systems, LLC, a Wisconsin Limited Liability Company of Brown Deer, Wisconsin is the real party in interest in this appeal.

II. RELATED APPEALS AND INTERFERENCES

There are no related appeals or interferences.

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III. STATUS OF CLAIMS

Claims 55-69 and 73-76 remain pending in this application and were finally rejected in the Office Action of July 8, 2002. All of these claims are on appeal. A copy of the appealed claims is contained in Appendix A. Claims and 1-54 and 70-72 have been cancelled and are not on appeal.

IV. STATUS OF AMENDMENTS

No Amendments after Final have been filed by Appellants.

V. SUMMARY OF THE INVENTION

Appellant's invention provides methods for identifying a DNA molecule. Specification page 1, lines 3-4. The methods involve hybridizing a single-stranded DNA molecule with a complementary reference strand of DNA to form a test duplex which is separated from at least one control duplex. Page 9, lines 9-14. The positions to which the test duplex and control duplex are detected, and the position of the test duplex is assigned an exact numerical migration value. Page 24, lines 23-32. The single-stranded DNA molecule is identified by matching the exact numerical migration value of the test duplex with a migration value in database containing the migration values of identified DNA molecules that is independent or separate from the separation. Page 24, lines 5-11 and 25-26, FIGS. 1 and 9-15.

VI. ISSUES

The issues on appeal are as follows:

1. Whether claims 55-69 and 73-76 meet the definiteness requirements of 35 U.S.C. § 112, second paragraph;
2. Whether claims 55-69 and 73-76 are nonobvious under 35 U.S.C. §103 over Zimmerman *et al.*, Nucleic Acids Research, Vol. 21, No. 19, 4541-4547 (1993), in view of Sapirstein *et al.*, Seed Science Technology 14, 489-517 (1986); and
3. Whether claims 55-69 and 73-76 are nonobvious under 35 U.S.C. §103 over Zimmerman *et al.*, in view of Sapirstein *et al.*, both further in view of Mullins *et al.*

VII. GROUPING OF THE CLAIMS

Claims 55-69 and 73-76 stand or fall together.

VIII. ARGUMENT

1. **The rejection under § 112, second paragraph, is clearly in error and should be withdrawn.**

In the Final Office Action dated July 8, 2002, claims 55-69 and 73-76 were “rejected under 35 U.S.C. 112, second paragraph, as being indefinite” for reciting “‘wherein the database of migration values is independent of the separation’ because it is not clear how this is possible.” Appellants respectfully disagree with this rejection and are unsure as to how using the word “independent” in accord with its common and ordinary meaning of “separate” or “apart from” renders the claims indefinite. *See, e.g.*, The Random House Thesaurus 373 (College ed. 1991), a copy of which is enclosed. In fact, the Examiner properly interpreted the meaning of this element stating, “[f]or the purposes of examination herein, the claims have been interpreted to mean that the database must be separate from the gel used for separation of the duplexes.” Because the claims apprise the skilled artisan of their meaning, Appellants respectfully request the rejection of claims 55-69 and 73-76 as indefinite be withdrawn.

As a point of clarification, Appellants also disagree with the Examiner’s statements that:

the database of migration values is intrinsically dependent upon the separation as the database is a record of the distances the test duplexes traveled in the separating step (b). Thus, it is not clear how the database can be independent from the separation. It seems that if the database were independent of the separation, the values contained in the database would be arbitrary values with no clear meaning or basis.

These statements appear to be based on a misunderstanding of the invention set forth in the claims. The present claims only recite the performance of a single separation – the separation between the test duplex and one or more control duplexes. No other separation is required to perform the claimed invention, although the performance of additional separations falls within the scope of the

claimed invention. The database is made up of “migration values of identified DNA molecules” but the claims do not require the separation of the identified DNA molecules or the calculation of their migration values be performed in the claimed separation of the test duplex from the one or more control duplexes. The migration values of the identified DNA molecules can be readily calculated from one or several separations that are not the claimed separation of the test duplex from the control duplex. As long as the migration values assigned in the different separations can be correlated to a common scale then the migration values can be easily related to one another. Thus, the database of values is not required to be “intrinsically dependent upon the (claimed) separation” or “a record of the distances the test duplexes traveled in the separating step (b).”

2. The claims are not obvious over Zimmerman *et al.* in view of Sapirstein *et al.*

In the Office Action of July 8, 2002, claims 55-59 and 73-76 were “rejected under 35 U.S.C. 103(a) as being unpatentable over Zimmerman *et al.* in view of Sapirstein *et al.* (Seed Science Technology (1986) 14(3) 489-517).” This rejection is improper not only because it is a hindsight reconstruction of the claimed invention, as Sapirstein *et al.* is not analogous to the claimed invention, but also because the references fail to teach or suggest all of the elements of the claimed invention. Moreover, the references are directed to different art areas and problems and teach away from their combination.

As discussed above, the present claims are directed to methods of identifying a DNA molecule using duplex separations and database matching. Zimmerman *et al.* teach genotyping DQA1 and DQB1 alleles using DNA probe hybridization and heteroduplex analysis. Contrast these to the non-analogous teachings of Sapirstein *et al.* which relate to wheat cultivar identification based on gliadin profiles. Sapirstein *et al.* do not disclose identification of anything at the molecular level, such as protein or DNA. Sapirstein *et al.* merely distinguish wheat cultivars. In order to discriminate between wheat cultivars, Sapirstein *et al.* do not use any hybridization technique, much less DNA hybridization. Moreover, Sapirstein *et al.* only care about protein banding patterns, not the identity of any specific protein in the banding pattern. In simplistic terms, if the banding pattern of sample X matches the database banding pattern for cultivar Y, then sample X is cultivar Y. The identity or sequence of the specific proteins that make up the banding

pattern are irrelevant. Accordingly, Appellants respectfully submit that a skilled artisan attempting to identify DNA would simply not look to art dealing with wheat cultivar banding for direction, other than in hindsight.

Additionally, the combination of Zimmerman *et al.* and Sapirstein *et al.* does not state a *prima facie* case of obviousness because they do not teach or suggest all of the elements of the appealed claims. Specifically, these references fail to teach or suggest at least two steps of the claimed invention, namely “assigning an exact numerical migration value to the position to which the test duplex migrates[,]” and “identifying the DNA molecule by matching the exact migration value with a database of migration values of identified DNA molecules, wherein the database of migration values is independent of the separation.”

The fact that the primary reference fails to teach or suggest these claim elements is explicitly admitted by the Examiner: “Zimmerman *et al.* do not assign an exact numerical migration value to the distance traveled by the heteroduplexes. Furthermore, Zimmerman *et al.* do not provide a database of migration values that is independent of (or separate from) the gel used in the separation step.” Despite the Examiner’s protests to the contrary, Zimmerman *et al.* also do not disclose a database of migration values of identified DNA molecules as required by the instant claims. To the skilled artisan a database is generally known as a “large collection of data organized especially for rapid search and retrieval (as by computer).” Merriam Webster’s Collegiate Dictionary 293 (Tenth Ed. 1997), a copy of which is enclosed. To meet this definition any database would intrinsically have to be separate and apart from a physical separation performed on heteroduplexes. Thus, the skilled artisan would not consider “[t]he left side of the gel in Figure 3...to be a database of test duplex migration values” as argued by the Examiner because the left side of the gel does not contain a “large collection of data organized especially for rapid search and retrieval” nor is the database separate and apart from the physical separation assay.

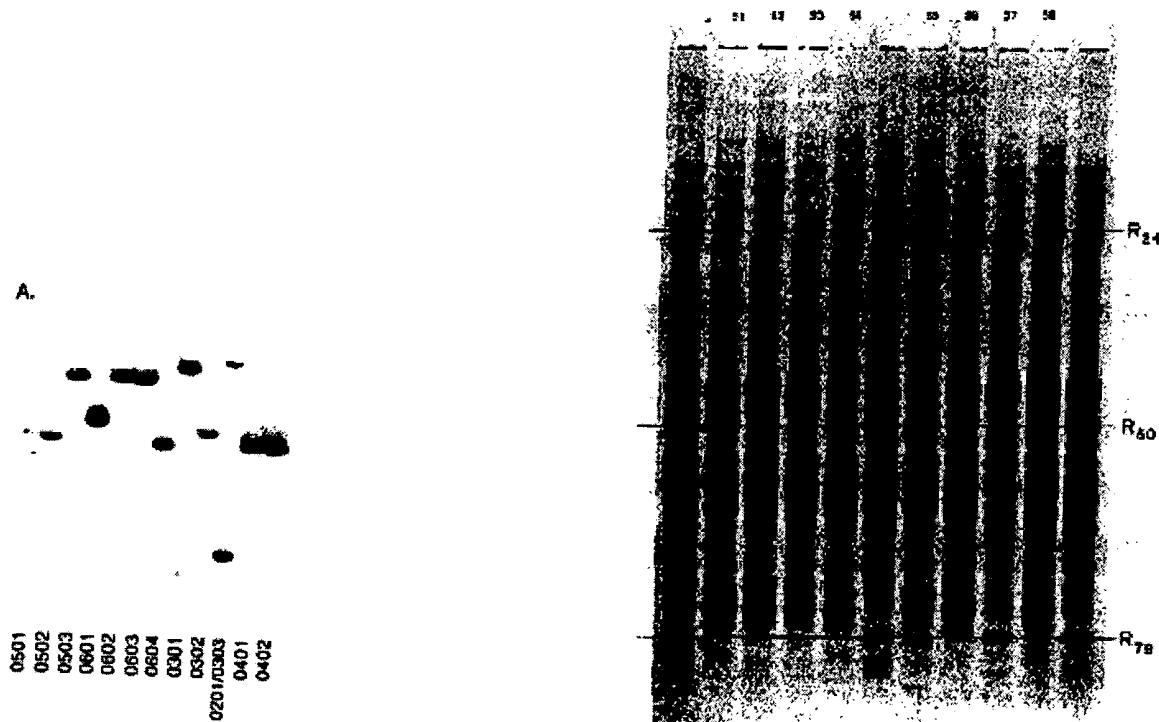
Instead, one skilled in the art would recognize the left side of the gel in figure 3 for what it is – a limited number of positive controls against which the heteroduplexes must be compared on the gel to identify the DNA molecules in the heteroduplex. In order to positively identify a truly unknown HLA allele in an individual, Zimmerman *et al.* would have to run well in excess of 100 positive controls. See Zimmerman *et al.* which note on page 4541 that the

HLA*DRB1, HLA*DQA1, HLA*DQB1, HLA*DPA1 and HLA*DPB1 genes contain 66, 10, 13, 4 and 26 separate alleles, respectively, for a grand total of 119 different alleles. These are only a fraction of the known HLA alleles which number in the hundreds, if not thousands. Schreuder *et al.*, *Tissue Antigens*, 58:109 (2001) (abstract enclosed). Therefore, Zimmerman *et al.* fails to teach several elements of the claimed invention.

Sapirstein *et al.* do not make up for the deficiencies of Zimmerman *et al.* because Sapirstein *et al.* also fail to teach or suggest these same elements of the claimed invention, namely “assigning an exact numerical migration value to the position to which the test duplex migrates; and identifying the DNA molecule by matching the exact migration value with a database of migration values of identified DNA molecules...” In fact, Sapirstein *et al.* deal only with individual proteins. Sapirstein *et al.* do not bother with hybridizing proteins to form duplexes, nor are they the least bit concerned with DNA. Although Sapirstein *et al.* disclose a database of gliadin protein band migration values nowhere do they disclose that any of the specific gliadin protein bands was identified or known. Sapirstein *et al.* only care about protein banding patterns, not the identity of any specific protein in the banding pattern. It is the combination of gliadin bands, not the individual bands themselves, that is important. The identity or sequence of the specific proteins that make up the banding pattern are irrelevant.

In order to overcome the deficiency of Zimmerman *et al.*, one skilled in that art would recognize that Sapirstein *et al.* would have to disclose a method for identifying a specific protein based on its position. This Sapirstein *et al.* do not disclose, instead their complete emphasis on banding patterns is evident throughout the article. See, e.g., Summary, p. 489 “unknown (or test) electropherograms (were compared) with counterpart protein patterns[.]” Therefore, Sapirstein *et al.* cannot teach or suggest “matching the exact migration value with a database of migration values of identified DNA molecules” recited in the claimed invention, and, as Zimmerman *et al.* also fail to teach this element, the combination of these two cannot render the claimed invention obvious. Instead, Appellants respectfully submit that the Examiner has cobbled together bits and pieces of non-analogous pieces of prior art in a hindsight reconstruction of the claimed invention.

Additionally, there is no suggestion or motivation in the references to combine their teaching. The Examiner forces together a combination of non-analogous references based not upon any teaching or suggestion in the cited references, but the advantages achieved by the claimed invention. At the same time the Examiner ignores the parts of the references that teach away from their combination. For example, Zimmerman *et al.* repeatedly point out that their method is simple and provides results that are easy to interpret. See, e.g., Page 4541, bottom of right column through page 4542, top of left column ("[t]his study presents a novel PCR-based approach... to achieve accurate and simple genotyping" and "this study has employed strategic labeling... to simplify HD banding patterns"). In contrast, Sapirstein *et al.* state that "standardising (protein) band migration distances... is sufficiently tedious to require implementation by means of a computer." Page 492, fourth paragraph. Contrast Zimmerman *et al.* where the "resulting HD pattern is comprised of a single product in homozygous individuals or two products in heterozygous individuals[,]" page 4542, top of left column, with Sapirstein *et al.* where the banding patterns consist of a large number of protein bands, an example of which is shown below:

Zimmerman *et al.*Sapirstein *et al.*

In fact, Sapirstein *et al.* reveal in Figures 5-8 that none of the numerous wheat cultivars studied provided a banding pattern that had less than 26 different bands. Figure 6, Pellisier cultivar.

Moreover, Sapirstein *et al.* rely on features not relevant to the teachings of Zimmerman *et al.*, such as complex statistical analysis and the subjective determination of “the intensity of the gliadin band(s)” as a “secondary feature” in the identification of wheat cultivars. Based on the divergent teachings of these references, the skilled artisan simply would not look to the complex, unrelated method of Sapirstein *et al.* for wheat cultivar identification to modify the simplified procedure of Zimmerman *et al.* for human lymphocyte typing.

In summary, not only is Sapirstein *et al.* not analogous to the claimed invention, the improper combination of Zimmerman *et al.* and Sapirstein *et al.* do not teach or suggest all elements of the claimed invention. Moreover, there is no suggestion or motivation other than hindsight to combine the references as suggested by the Examiner. In reality, a careful reading of the references reveals they teach away from their combination. Therefore, the rejection of the claims as being obvious over Zimmerman *et al.* and Sapirstein *et al.* should be withdrawn.

3. The claims are not obvious over Zimmerman *et al.* and Sapirstein *et al.* both in view of Mullins *et al.*

Claims 55-69 and 73-76 were also “rejected under 35 U.S.C. 103(a) as being unpatentable over Zimmerman *et al.* in view of Sapirstein *et al.* (Seed Science Technology (1986) 14(3) 489-517) both further in view of Mullins *et al.*.” The teachings of Zimmerman *et al.* and Sapirstein *et al.* are discussed above. Mullins *et al.* were cited by the Examiner “merely to demonstrate that it was known in the art at the time the invention was made to determine exact numerical mobility values for heteroduplex nucleic acid molecules.”

However, the teachings of Mullins *et al.* are irrelevant to the claimed invention because they do not calculate the exact migration distance of a single DNA duplex as the Examiner contends. Instead, in example 4 of Mullins *et al.* cited by the Examiner, migration values “were calculated as the average distance of migration of the two heteroduplex bands divided by the distance of migration of the homoduplex bands.” Page 50, line 37 through page 51, line 2. Because the migration values are based on an “average distance of two heteroduplex bands” the

migration value relates to both of these heteroduplexes and is not specific for either one of them. Thus, any migration value calculated in Mullins *et al.* is simply not capable of providing for the positive and actual identification of any of the DNA molecules in the heteroduplex bands as is required by the claimed invention. Moreover, because Mullins *et al.* are cited solely for the proposition that mobility values of heteroduplexes can be measured this reference must also fail to overcome the deficiencies of Zimmerman *et al.* and Sapirstein *et al.* discussed above. Therefore, the rejection of the claims as being obvious over Zimmerman *et al.* and Sapirstein *et al.* both in view of Mullins *et al.* should be withdrawn.

IX. CONCLUSION

For the foregoing reasons, Appellants submit that Claims 55-69 and 73-76 are patentable over the cited references, and that the final rejection of these claims should be reversed.

Respectfully submitted,

Date December 6, 2002

By 

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APPENDIX A - CLAIMS ON APPEAL

55. A method for identifying a DNA molecule comprising:

(a) hybridizing a single strand DNA molecule with a complementary reference DNA strand to form a test duplex;

(b) separating the test duplex from at least one control duplex run in the same separation;

(c) detecting the positions to which the test duplex and the at least one control duplex migrate in the separation;

(d) assigning an exact numerical migration value to the position to which the test duplex migrates; and

(e) identifying the DNA molecule by matching the exact migration value with a database of migration values of identified DNA molecules, wherein the database of migration values is independent of the separation.

56. The method of claim 55 wherein the complementary reference DNA strand is labeled.

57. The method of claim 55 further comprising repeating steps (a)—(e) one or more times wherein a different complementary reference strand is utilized in each repeat of steps (a)-(e) to identify the DNA molecule.

58. The method of claim 55 wherein step (b) comprises gel electrophoresis.

59. The method of claim 55 wherein the database of migration values comprises migration values of alleles of a gene selected from the group consisting of HLA, TAP, LMP, ras, non-classical HLA and Bf.

60. The method of claim 55 wherein the database of migration values comprises migration values for mammalian MHC genes.

61. The method of claim 59 wherein the alleles are selected from the group consisting of HLA alleles.

62. The method of claim 60 wherein the DNA molecule comprises a portion of an HLA gene.

63. The method of claim 55 further comprising confirming the identity of the DNA molecule by sequencing the test duplex, performing sequence specific primer amplification analysis on the test duplex or performing sequence specific oligonucleotide analysis on the test duplex.

64. The method of claim 55 wherein the method can resolve a difference of one, two or three nucleic acid positions between the DNA molecule and the complementary sequence of the complementary reference DNA strand.

65. The method of claim 55 wherein the DNA molecule and complementary reference DNA strand have the same number of nucleotides.

66. The method of claim 55 wherein the complementary reference DNA strand consists of the wild type sequence of a naturally occurring DNA strand of interest or a naturally occurring mutant thereof.

67. The method of claim 55 wherein the at least one control duplex comprises (i) duplexes which have faster and slower mobility than the test duplex or (ii) duplexes which have graded mobilities.

68. The method of claim 55 wherein the identified DNA molecule is matched to a second identified DNA molecule and the method is used to match tissue between a prospective tissue donor and a prospective tissue recipient.

69. The method of claim 55 further comprising amplifying the DNA molecule prior to step (a).

73. A method for identifying a DNA molecule, comprising:

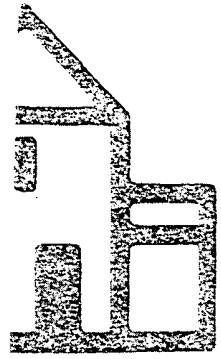
- (a) amplifying a DNA molecule to produce amplified double stranded DNA molecules;
- (b) denaturing the amplified double stranded DNA molecules into single stranded DNA molecules wherein the single stranded DNA molecules include sense and antisense strands;
- (c) hybridizing the single stranded DNA molecules with reference DNA strands which are complementary to the single stranded DNA molecules to form test duplexes;
- (d) separating the test duplexes from at least one control duplex run in the same separation;
- (e) detecting the positions to which the test duplexes and the at least one control duplex migrate in the separation;
- (f) assigning an exact numerical migration value to the position to which the test duplex migrates; and
- (g) identifying the DNA molecule by matching the exact migration with a database of migration values of identified DNA molecules, wherein the database of migration values is independent of the separation.

74. The method of claim 73 wherein the reference DNA strands are labeled.

75. The method of claim 73 further comprising repeating steps (a)-(g) one or more times wherein a different complementary reference strand is utilized in each repeat of steps (a)-(g) to identify the DNA molecule.

76. The method of claim 73 wherein step (f) comprises:

- (i) assigning a migration value to the at least one control duplex; and
- (ii) assigning the exact numerical migration value to the test duplex based on the relative migration position of the test duplex compared to the migration value of the control duplex.



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indentation

apart, as a matter of fact, to tell honest, in point of fact, actually, vñ to it.

j. The new man is an indefatigable and will go far: tireless, inexhausting, persistent, diligent, c, sedulous, unfaltering, unflag-wearying, staunch.

i. 1 The indefensible village fell hours: unable to withstand defenseless; open to attack, untenable, vincible. 2 Her purchase was an indefensible waste of bale, beyond justification, unjustifyable, unpardonable, without due; open to criticism, improper,

e, invincible, invulnerable, probable, justifiable, excusable, parable, rational, tenable.

The store will be closed for an d: unspecified, having no fixed ined, indeterminate, unknown, it; illimitable, measureless, limm an indefinite answer unse indistinct, vague; not clearly defined, amorphous, indecisive, ve, obscure, ambiguous, unsure.

e, clear-cut, certain, limited,

2 definite, certain, settled, sure,

The book left an indelible ne. The ink made an indelible rmanent, unforgettable, lasting manently fixed, fast, ineradicab; unremoveable, incapable of or wiped out, ingrained, deep-

His indelicate manners are the tigh upbringing: coarse, crude, unbecoming; clumsy, awkward. ndelicate story to tell in mixed sive, lacking good taste, indis- unrefined; off-color, immodest, vulgar, suggestive, indecorous, oad, obscene, coarse, gross, im-

e, refined, seemly, decorous, polished. 2 decent, chaste, proper.

be company indemnified him in his car: reimburse, repay, pay ite, requite, remunerate, recomp ht, make restitution, make up l, rectify, make amends, satisfy,

1 You can identify some trees tions of the leaves. The map us indentations along the shore- l, incision, cavity, furrow, score; ity, niche, pocket, bay, inset. 2

independence

High heels made indentations in the linoleum: dent, pit, gouge, depression, nick.

Ant. bump, protuberance, projection, prominence.

independence *n. The American colonies won their independence from England: emancipation, liberty, freedom, self-determination, self-government, freedom from control, sovereignty, self-reliance, liberation, autonomy.*

Ant. dependence; subordination, subjection; servitude, slavery, bondage; reliance, dependency.

independent *adj. 1 Children should be encouraged to be independent thinkers: self-reliant, uncontrolled, on one's own, autonomous, free, self-directing, individualistic, uncoerced; unconstrained, free from the control or influence of others. 2 The United States became an independent country in 1776: free, self-governing, autonomous; self-determining, sovereign. 3 The medical college is independent of the university: separate, not joined to or associated with, unattached to, distinct from, exclusive, apart from, unconnected with, unallied. 4 The inheritance made him independent for life: solvent, well-to-do, well-off, in easy circumstances, well-fixed, affluent, Informal well-heeled.*

Ant. 1 dependent; influenced, controlled, directed; subordinate, subject, tributary, subservient, servile, slavish, subject, attached, interrelated. 4 dependent, reliant, beholden, attached.

indescribable *adj. The artistry of Michelangelo's Pietà is indescribable: beyond description, inexpressible, beyond words, ineffable, beggar description; indefinable; overwhelming, unutterable.*

indestructible *adj. Toys for toddlers should be as indestructible as possible: unbreakable, damage-resistant, infrangible, enduring, permanent, everlasting, imperishable, incapable of being destroyed.*

Ant. destructible, perishable, breakable, unenduring, fragile, frangible.

indeterminate *adj. The acting superintendent will serve for an indeterminate time: unspecified, undetermined, unstipulated, uncertain, unfixed in extent or amount, unclear, obscure, not clear, unresolved, vague, undefined; ambiguous, problematic, perplexing, indefinite.*

Ant. specified, precise, definite, clear, certain, determined, fixed, defined.

index *n. 1 The poems are given in the index by author: alphabetical list, catalog, register, glossary. 2 Overeating is often an index of emotional stress: sign, token, indication, indicator, symptom, clue, evidence, manifestation, proof, mark.*

indicate *v. 1 The patient's pallor indicates anemia: be a sign of, be symptomatic of, show, designate, denote, imply, point to, suggest; evince, bespeak, reveal, symbolize, signify, stand for, mean, represent. 2 Indicate where the pain is*

indigent

with your finger: point out, point to, specify, direct attention to. 3 A barometer indicates air pressure: show, make known, register, reveal, tell, establish, record.

indication *n. There was no indication this morning that it would rain: sign, hint, intimation, signal, manifestation, token, warning, evidence, mark, clue, suggestion, foretoken, hint, symptom, gesture, demonstration; portent, augury, omen, boding, foreboding, premonition, presage; signifying, telling, indicating, designation, mention, showing, pointing.*

indicative *adj. Fever may be indicative of infection: suggestive, indicatory, characteristic, evidential, symptomatical, symptomatic, expressive, significant, emblematic, symbolic, representative, denotative, connotative, designative.*

indict *v. The grand jury indicted him for embezzlement: arraign, accuse, charge, inculpate, criminate, impute, bring to justice, cite, impeach, prosecute; find an indictment against, prefer charges, have up, pull up, bring up.*

indifference *n. 1 He would have preferred anger to her cold indifference: unconcern, absence of feeling, lack of interest, disinterest, neglect, inattention, impassiveness, impassivity, nonchalance, aloofness, carelessness, negligence, insensibility, insensitivity, disdain, insouciance, apathy, coldness. 2 Religion is a matter of indifference to many of today's youth: unimportance, triviality, no import, insignificance, paltriness, inconsiderableness.*

Ant. 1 concern, warmth, sensibility; attention, interest, eagerness, caring. 2 importance, magnitude, significance, greatness.

indifferent *adj. 1 The writer was indifferent to criticism, good or bad: unconcerned, not caring, insensible, without interest, impervious, uninterested, insusceptible, detached, unmindful, impassive, unmoved, insouciant, apathetic, nonchalant, cool, aloof. 2 The actor's indifferent performance left the audience cold: perfunctory, mediocre, not very good, undistinguished, uninspired, ordinary, so-so, rote, commonplace, neither good nor bad, medium, middling, fair, modest, moderate, passable, average, betwixt and between; falling short of any standard of excellence, second-rate, rather poor.*

Ant. 1 avid, eager, keen, agog; interested, sensitive, susceptible, caring, sympathetic, responsive, compassionate, enthusiastic. 2 choice, notable, remarkable, exceptional, rare, first-class, superior, excellent, fine.

indigenous *adj. Cotton is indigenous to the southern United States: native, growing naturally, aboriginal, originating in, characteristic of, endemic, homebred, home-grown, domestic, autochthonous.*

Ant. naturalized; exotic; foreign, alien, extraneous, imported.

indigent *adj. Many indigent people receive government aid: needy, destitute, in want, lack-*

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Abstract

Tissue Antigens
Volume 58 Issue 2 Page 109 - August 2001

The HLA Dictionary 2001: a summary of HLAA, -B, -C, -DRB1/3/4/5, -DQB1 alleles and their association with serologically defined HLA-A, -B, -C, -DR and -DQ antigens

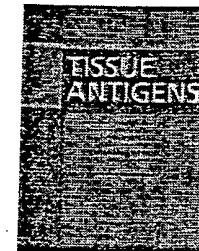
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Abstract: This report presents the serologic equivalents of 123 HLA-A, 272 HLA-B and 155 HLA-DRB1 alleles. The equivalents cover over 64% of the presently identified HLA-A, -B and -DRB1 alleles. The dictionary is an update of the one published in 1999 and also includes equivalents for HLA-C, DRB3, DRB4, DRB5 and DQB1 alleles. The data summarize information obtained by the WHO Nomenclature Committee for Factors of the HLA System, the International Cell Exchange (UCLA), the National Marrow Donor Program (NMDP) and individual laboratories. In addition, a listing is provided of alleles which are expressed as antigens with serologic reaction patterns that differ from the well-established HLA specificities. The equivalents provided will be useful in guiding searches for unrelated hematopoietic stem cell donors in which patients and/or potential donors are typed by either serology or DNA-based methods. These equivalents will also serve typing and matching procedures for organ transplant programs where HLA typings from donors and from recipients on waiting lists represent mixtures of serologic and molecular typings. The tables with HLA equivalents and a questionnaire for submission of serologic reaction patterns for poorly identified allelic products will also be available on the WMDA web page: <http://www.worldmarrow.org>.

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- antigens
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World Marrow Donor Association
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